5

10

MGRGDGAGAFGLGFTPPHGGLLGWSPQAQGILETLPANPPPASTNRQSGRQPTPLSP PLRNTHPQAMQWNSTTFHQTLQDPRVRGLYFPAGGSSSGTVNPVPTTVSPISSIFSRIG DPALNMENITSGFLGPLLVLQAGFFLLTRILTIPQSLDSWWTSLNFLGGTTBCLGQNS QSPTSNHSPTSCPPTCPGYRWMCLRRFIIFLFILLLCLIFLLVLLDYQGMLPVCPLIPGSS TTSTGPCRTCTTPAQGTSMYPSCCCTKPSDGNCTCIPIPSSWAFGKFLWEWASARFS WLSLLVPFVQWFVGLSPTVWLSVIWMMWYWGPSLYSILSPFLPLLPIFFCLWVYI (SEQ. ID NO: _____)

FIG. 1

MPLSSIFSRIGDPTVQASKLCLGWLWGMDIDPYKEFGATVELLSFLPSDFFPSVRDLL

10 DTASALYREALESPEHCSPHHTALRQAILCWGELMTLATWVGVNLEDPEF<u>RGD</u>ASR
DLVVSYVNTNMGLKFRQLLWFHISCLTFGRETVIEYVLSFGVWIRTPPAYRPPNAPIL

STLPETTVVRRRGRSPRRRTPSPRRRRSQSPRRRRSQSREPQC

FIG. 2